TODD GARNER DS 6306 WEEK 6 PART 1 FEBRUARY 87. 2023

READ IN THE DATA AND CLEAN - PART 1, #1

• The site www.opendatasoft.com proved challenging so I found the .csv files on kaggle.com and also in the class GitHub repo.

```
96 ```{r}
97  # Load the Titanic dataset
98  Titanic <- read.csv(file.choose(), header = TRUE)
99  library(dplyr)
100  Titanic_new <- Titanic %>% select(Survived, Pclass, Age)
101  Titanic_exp <- Titanic %>% select(Pclass, Age)
102  Titanic_new_final <- Titanic_new[complete.cases(Titanic_new), ]
103  Titanic_final <- Titanic_exp[complete.cases(Titanic_exp), ]|
```

- Two data sets: training and test. Removed the NA's and selected the three pertinent columns.
- Comment: I used ChatGPT and found it distracted me more than it assisted me. I may use it in a different manner than writing the code base.

CREATE THE TRAINING/TESTING SETS, USE THEM AND CONFUSION MATRIX – PART 1, #2

```
115 set.seed(123)
116 train_new_index = sample(1:nrow(Titanic_new_final), size = nrow(Titanic_new_final)*0.8, replace = FALSE)
117 train_new = Titanic_new_final[train_new_index, ]
118 test_new = Titanic_new_final[-train_new_index, ]
119
120 set.seed(123)
121 train_index = sample(1:nrow(Titanic_final), size = nrow(Titanic_final)*0.8, replace = FALSE)
122 train = Titanic_final[train_index,
123 test = Titanic_final[-train_index,
124
125 library(class)
126 library(caret)
127 knn.23 <- knn(train, test, train_new$Survived, k = 23)
128 knn. 24 <- knn(train, test, train_new$Survived, k = 24)
129 table(test_new$Survived, knn.23)
     confusionMatrix(table(test_new$Survived, knn.23))
131
132 table(test_new$Survived, knn.24)
     confusionMatrix(table(test_new$Survived, knn.24))
```

- Data set: training -> 571 testing -> 143 (177 NA's removed)
- Transforming in KNN and Confusion Matrix (results on next slide)

CONFUSION MATRIX AND STATISTICS

```
Confusion Matrix and Statistics
  knn. 24
    0 1
 0 69 10
 1 46 18
              Accuracy: 0.6084
                95% CI: (0.5233, 0.6889)
   No Information Rate: 0.8042
   P-Value [Acc > NIR] : 1
                 Kappa: 0.1634
Mcnemar's Test P-Value : 2.91e-06
           Sensitivity: 0.6000
           Specificity: 0.6429
        Pos Pred Value: 0.8734
        Neg Pred Value: 0.2813
            Prevalence: 0.8042
        Detection Rate: 0.4825
  Detection Prevalence: 0.5524
     Balanced Accuracy: 0.6214
       'Positive' Class: 0
```

Having seen the answers, I know now that my solution was off the mark. I'm impressed with your code. Specific and on target. I have much to learn.

AGE AND PROBABILITY OF SURVIVAL, PART 1, #3

Through my research, I found one suggestion on the value to select for k. It was suggested that k can be derived from the square root of the size of the data set.

The result is between k = 23 and k = 24. I've employed them both

I created 3 data sets with my Age (59) in it along with the Pclass. 1st, 2nd, 3rd.

Given the data, it looks bad for someone my age, regardless of the class.

[1] 0
Levels: 0 1
[1] 0
Levels: 0 1
[1] 0
Levels: 0 1

USE OF THE TRAINING SET AND THE TEST SET – PART 1 #4

• Utilizing the prior training set (571 observations), I've loaded in the test_set.csv. This is where I got stuck and moved to Part 2.

```
165 * ```{r}
166  Titanic_418 <- read.csv(file.choose(), header = TRUE)
167  Titanic_418_new <- Titanic_418 %>% select(Pclass, Age)
168  Titanic_418_final <- Titanic_418_new[complete.cases(Titanic_418_new), ]
169  knn.418_final <- knn(train, Titanic_418_final, train_new$Survived, k = 18)
170  train_new$Survived <- as.factor(train_new$Survived)
```

PART 2 # A. IRIS DATA SET – 70/30 TRAINING SET, PLOT XAXIS

```
19 library(tidyverse)
20 iris
21 View(iris)
22 splitPerc = .70
23 train_init = sample(1:dim(iris)[1],round(splitPerc * dim(iris)[1]))
24 train = iris[train_init,]
25 test = iris[-train_init,]
26 . . . .
       Description: df [150 x 5]
                        Sepal.Length
<dbl>
                                                                                    Petal.Length
<dbl>
                                                      Sepal.Width

«dbl»
                                                                                                                  Petal.Width Species
                                                                                                                          0.2 setosa
                                  5.1
                                                              3.5
                                                                                             1.4
                                                                                             1.4
                                                                                                                          0.2 setosa
                                  4.9
                                                               3.0
                                                                                             1.3
                                                                                                                          0.2 setosa
                                  4.7
                                                              3.2
                                                                                             1.5
                                  4.6
                                                              3.1
                                                                                                                          0.2 setosa
                                  5.0
                                                              3.6
                                                                                             1.4
                                                                                                                          0.2 setosa
                                  5.4
                                                                                             1.7
                                                              3.9
                                                                                                                          0.4 setosa
                                  4.6
                                                                                             1.4
                                                              3.4
                                                                                                                          0.3 setosa
                                                                                             1.5
                                  5.0
                                                              3.4
                                                                                                                          0.2 setosa
                                                                                             1.4
                                  4.4
                                                              2.9
                                                                                                                          0.2 setosa
                                                                                             1.5
                                                                                                                          0.1 setosa
                                  4.9
                                                              3.1
```

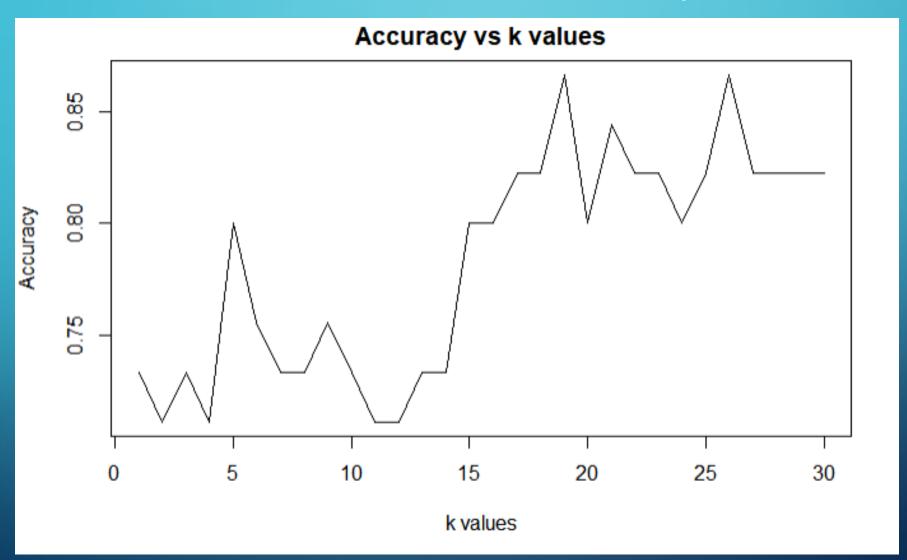
>70/30 TRAIN/TEST | ITERATE ALL K VALUES

```
32 iris
33 library(class)
34 library(FNN)
35 library(caret)
36 library(ISLR)
37 set.seed(6)
38 splitPerc = .70
39 trainIndices = sample(1:dim(iris)[1],round(splitPerc * dim(iris)[1]))
40 train = iris[trainIndices,]
41 test = iris[-trainIndices,]
42 iris %>% ggplot(aes(x = Sepal.Length,Sepal.Width,color = Species)) + geom_point()
43 View(train$Species)
44 # k = 5
  classifications = knn(train[,c(1,2)],test[,c(1,2)],train$Species, prob = TRUE, k = 5)
46 table(test$Species,classifications)
47 confusionMatrix(table(test\Species.classifications)\)
49 accs = data.frame(accuracy = numeric(30), k = numeric(30))
50 nrow(train)
51 for(i in 1:30)
52 √ {
  classifications = knn(train[,c(1,2)],test[,c(1,2)],train\\ species, prob = TRUE, k = i)
54 table(test$Species,classifications)
  CM = confusionMatrix(table(test$Species,classifications))
  accs$accuracy[i] = CM$overall[1]
58 △ }
59 plot(accs$k,accs$accuracy, type = "l")
```

Iris – Split the data into 70/30 train/test sets, plot the results (plots on the following slide), classify and employ the confusion matrix.

Create a "for loop" to test the k values to find the best possible fit.

DETERMINATION OF BEST K VALUE | ANSWER = 18



CONFUSION MATRIX RESULTS

```
13
  setosa
                            0
                                      0
 versicolor
                 1
                           11
 virginica
                                     13
                 0
                            3
Overall Statistics
              Accuracy: 0.8222
                95% CI: (0.6795, 0.92)
   No Information Rate: 0.3778
   P-Value [Acc > NIR] : 1.262e-09
                 Карра: 0.7327
Mcnemar's Test P-Value: NA
Statistics by Class:
                    Class: setosa Class: versicolor Class: virginica
Sensitivity
                           0.9286
                                             0.7857
                                                             0.7647
Specificity
                           1.0000
                                             0.8387
                                                             0.8929
Pos Pred Value
                           1.0000
                                             0.6875
                                                             0.8125
Neg Pred Value
                           0.9688
                                             0.8966
                                                             0.8621
Prevalence
                           0.3111
                                             0.3111
                                                             0.3778
                           0.2889
                                             0.2444
Detection Rate
                                                             0.2889
Detection Prevalence
                           0.2889
                                             0.3556
                                                             0.3556
Balanced Accuracy
                           0.9643
                                             0.8122
                                                             0.8288
[1] 105
```

BONUS QUESTION: REPEAT PRIOR ANALYSIS WITH LOOCV METHODOLOGY - RESULTS

setosa	50	0	0			
versicolor	0	31	12			
virginica	0	19	38			
Overall Statistics						
95% No Information R P-Value [Acc > N	CI: ate: IR]: ppa:	< 2.2e-16	. 8551)			
1						
	class	: setosa C	lass: vers	sicolor	class:	virginica
Sensitivity		1.0000		0.6200		0.7600
Specificity		1.0000		0.8800		0.8100
Pos Pred Value		1.0000		0.7209		0.6667
Neg Pred Value		1.0000		0.8224		0.8710
Prevalence		0.3333		0.3333		0.3333
Detection Rate		0.3333		0.2067		0.2533
Detection Prevalence		0.3333		0.2867		0.3800
Balanced Accuracy		1.0000		0.7500		0.7850
[1] 105						

These results are lower slightly than the k=18 confusion matrix results on the prior page.

I think this highlights that LOOCV is great to use on small-ish data sets. On data sets with hundreds of values/observations, it appears that LOOCV is not superior to determining an external (XX%/XX%) versus an internal (LOOCV) methodology.